

CLN00539416.a	MLTLGEQLPLVTRFRHSVHPPRLDLFFIWIWFCFYVFLKNNFLGNLLDYNCS SSPMQRKG	60
CLN00250082.b	MLTLGEQLPLVTRFRHSVHPPRLDLFFIWIWFCFYVFLKNNFLGNLLDYNCS SSPMQRKG	60
24980850_24980849	-----	0
CLN00539416.a	SYCWDFEEMVLEQLEIHTHTKNLNPYLTPDTKATFK-----	96
CLN00250082.b	SYCWDFEEAVRCHWAV-----	76
24980850_24980849	-----MEPSGSEQLFEDPDPGGKSQDAEARK	26
CLN00539416.a	-----	96
CLN00250082.b	-----	76
24980850_24980849	QTESEQKLSKMTNNALENINVIGQGLKHLFQHQRSSSVSPHDVQQIQADPEPEMDLESQ	86
CLN00539416.a	-----	96
CLN00250082.b	-----	76
24980850_24980849	NACAEIDGVPTHTPTALNRVLQQIRVPPKMKRGTSLSHRRGKPEAPKGSPOINRKSGQEMT	146
CLN00539416.a	-----	96
CLN00250082.b	-----	76
24980850_24980849	AVMQSGRPRSSSTTDAPTGSAMMEIACAAAAAAAACLPGEETAERIERLEVSSLAQTSS	206
CLN00539416.a	-----	96
CLN00250082.b	-----	76
24980850_24980849	AVASSTDGSIHTDSVDGTPDPTQRTKAAIAHLQQKILKLTQIKIAQTARDDNVAEYKLA	266
CLN00539416.a	-----	96
CLN00250082.b	-----	76
24980850_24980849	NSADKQQAARIKQVFEKKNQKSAQTILQLQKLEHYHRKLREVEQNGIPROPKDVFRDMH	326
CLN00539416.a	-----	96
CLN00250082.b	-----	76
24980850_24980849	QGLKDVGAQVTGTFSEGVVDSVKGGFSSFSQATHSAAGAVVSKPREIASLIRNKFGSADNI	386
CLN00539416.a	-----	96
CLN00250082.b	-----	76
24980850_24980849	PNLKDSLEEGQVDDAGKALGVISNFQSSPKYGSEEDCSSATSGSVGANSTTGGIAGVASS	446
CLN00539416.a	-----	96
CLN00250082.b	-----	76
24980850_24980849	SKTNTLDMQSSGFDALLHEIQEIRETQARLEESFETLKEHYQRDYSLIMQTLQEERYRCE	506
CLN00539416.a	-----	96
CLN00250082.b	-----	76
24980850_24980849	RLEEQLNDLTTELHONEILNLKQELASMEEKIAYQSYERARDIQEALACQTRISKMELQQ	566
CLN00539416.a	-----	96
CLN00250082.b	-----	76
24980850_24980849	QQQVVVQLEGLLENATARNLLGKLINILLAVMAVLLVFVSTVANCVVPLMKTRNRTFSTLF	626
CLN00539416.a	-----	96
CLN00250082.b	-----	76
24980850_24980849	LVVFI AFLWKHWDALFSYVERFFSSPR	653

FIG. 1

CLN00219153.a	-----
24980850_24980849	MEPSGSEQLFEDPDPPGGKSQDAEARKQTESEQKLSKMTNNALENINVIGQGLKHLFQHQH
CLN00149041.a	MEPSGSEQLFEDPDPPGGKSQDAEARKQTESEQKLSKMTNNALENINVIGQGLKHLFQHQH
CLN00219153.a	-----
24980850_24980849	RRSSVSPHDVQQIQADPEPEMDLESQNAEIDGVPTHPTALNRVLQQIRVPPKMKRGTS
CLN00149041.a	RRSSVSPHDVQQIQADPEPEMDLESQNAEIDGVPTHPTALNRVLQQIRVPPKMKRGTS
CLN00219153.a	-----
24980850_24980849	LHSRRGKPEAPKGSPOINRKSGQEMTAVMQSGRPRSSSTTDAPTGSAMMEIACAAAAAA
CLN00149041.a	LHSRRGKPEAPKGSPOINRKSGQEMTAVMQSGRPRSSSTTDAPTSSAMMEIACAAAAAA
CLN00219153.a	-----
24980850_24980849	ACLPGEEGTAEIERLERLEVSSLAQTSSAVASSTDGSIHTDSVDGTPDPQRTKAAIAHLQOK
CLN00149041.a	ACLPGEMPLPGFTWANSSTS-----
CLN00219153.a	-----
24980850_24980849	ILKLTEQIKIAQTARDDNVAEYLLANSADKQQAARIKQVFEKKNQKSAOTILOLQKLE
CLN00149041.a	-----
CLN00219153.a	-----
24980850_24980849	HYHRKLREVEQNGIPROPKDVFRDMHQGLKDVGAQVTGFSQGVVDSVKGGFSSFSQATHS
CLN00149041.a	-----
CLN00219153.a	-----
24980850_24980849	AAGAVVSKPREIASLIRNKFGSADNIPNLKDSLEEGQVDDAGKALGVISNPFQSSPKYGSE
CLN00149041.a	-----
CLN00219153.a	-----
24980850_24980849	EDCSSATSGSVGANSTTGGIAVGASSSKTNTLDMQSSGFDALLHEIQEIRETOARLEESF
CLN00149041.a	-----
CLN00219153.a	-----
24980850_24980849	ETLKEHYQRDYSLLIMQTLQEERYRCERLEEQLNDLTELHONEILNLKQELASMEEKIAYQ
CLN00149041.a	-----
CLN00219153.a	-----
24980850_24980849	SYERARDIQEAEACQTRISKMELOQQQQQVVOLEGLLENATARNLLGKLINILLAVMAVL
CLN00149041.a	SYERARDIQEAEACQTRISKMELOQQQQQVVOLEGLLENATARNLLGKLINILLAVMAVL
CLN00219153.a	-----
24980850_24980849	LVFVSTVANCVVPLMKTRNRTFSTLFLVVFIAFLWKHWDALFSYVERFFSSPR 121
CLN00149041.a	LVFVSTVANCVVPLMKTRNRTFSTLFLVVFIAFLWKHWDALFSYVERFFSSPR 653
	-----SWLSWQSFWSLSPL 214
	: : * : : **

FIG. 2

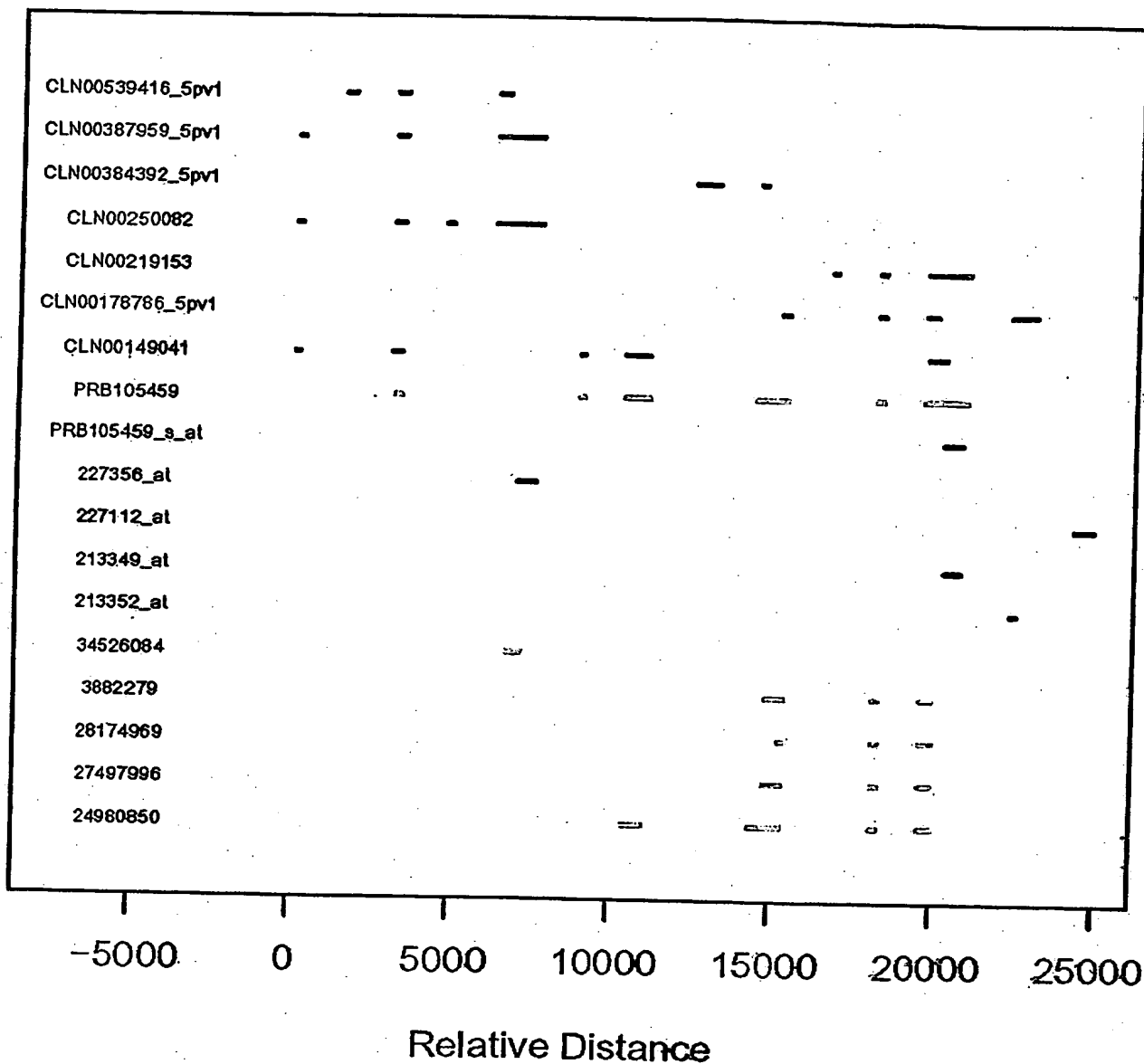


FIG. 3

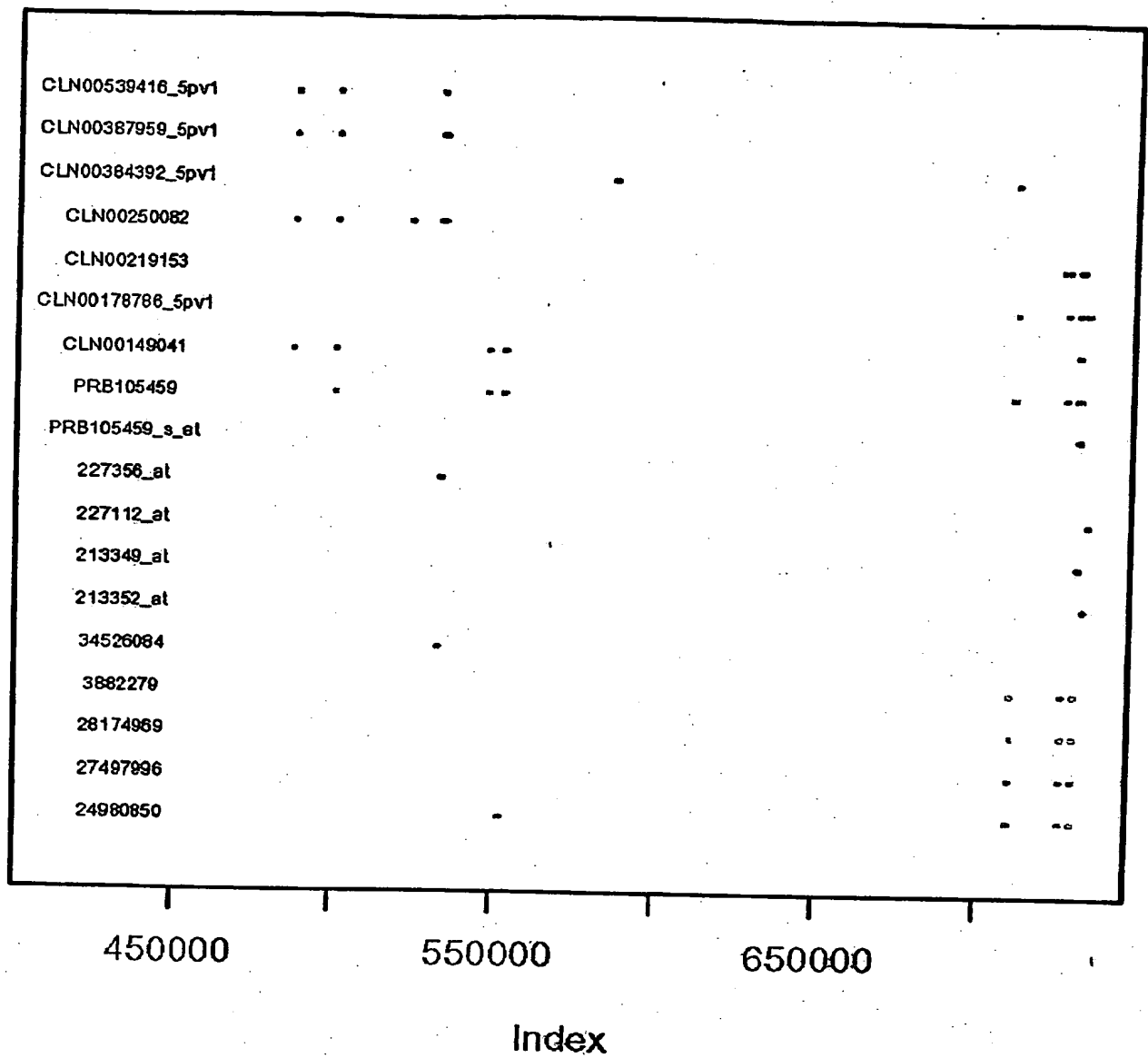


FIG. 4

Best Available Copy